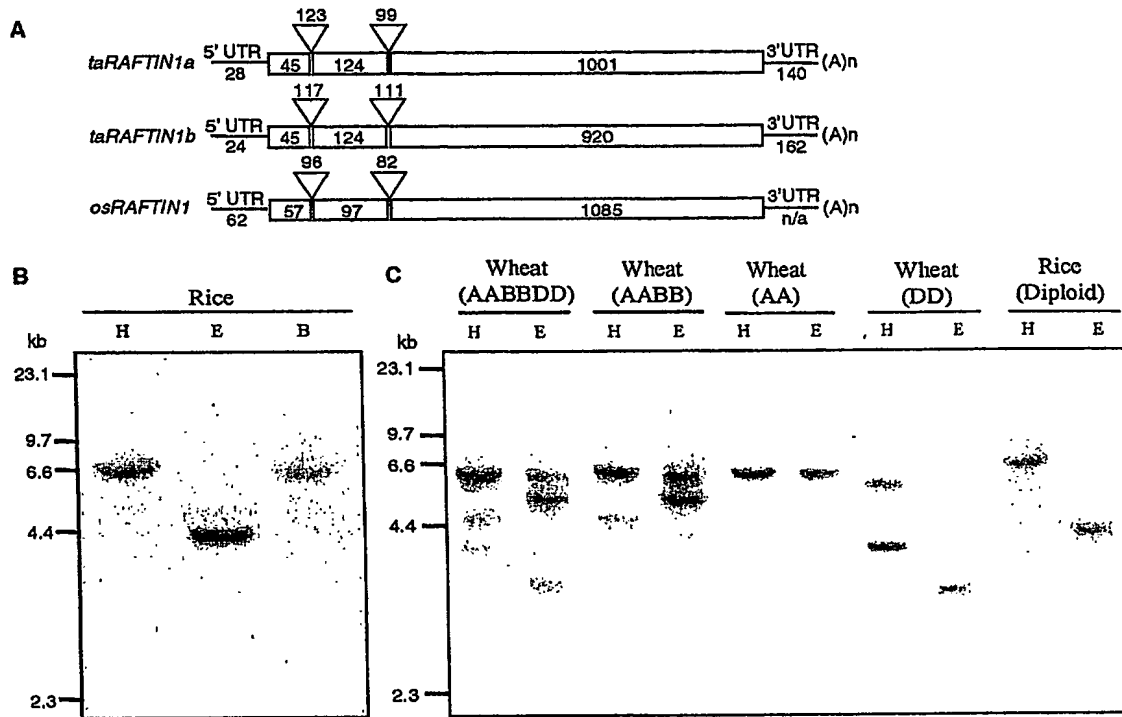
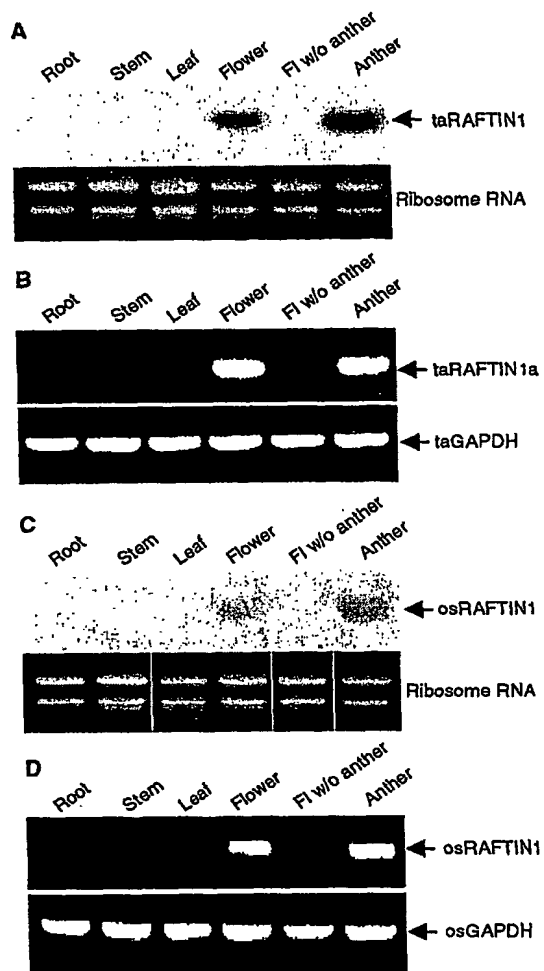


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Figure 2 Aiming Wang *et al.* 2002Top
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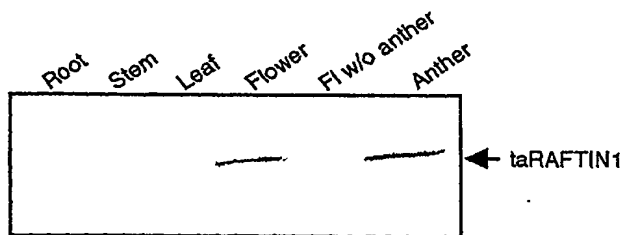


Figure 3 Aiming Wang *et al.* 2002



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A

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taRAFTIN1a 1  MARFLVALLATTILVAVOAGGQLGHAAPATAEVEFWRVLPSPPLDAVLRLLLKQPAAGVELLTEATSFVRDAEDRAAPFD 78
taRAFTIN1b 1  -----G-----S-----P-----AA----- 72
osRAFTIN1 1  -----AA--LVAVA-AA-VLS--D--S-----F-----RPDTSF-VGKA--ACAAAAGAARTGF-- 74

taRAFTIN1a 79  YRDYSRSPDDEPSKSTGAAASGARDPDYDDYSAAAAGGDKLRGAASGARAAAAADFDYDDYSGADKLRGATDAAAA 141
taRAFTIN1b 73  -----SS-----V-AA-----AAAAAAAAAAAAAA-----N-----ER-----AAAA 114
osRAFTIN1 75  -T--RG-DSPTTA-GLDL-GDFGEFAP-G-A--AQGE--GGGAA--A-EQVLAVDAG-N--K-V--R-L--GSSTAGGE 153

taRAFTIN1a 142  AAAAAAAEYKAPSSSLAGNGASMARGAGKAETTTVFHEEAVRVGKRLPFRFPPTPAALGFLPRQVADSVPTTAALP 212
taRAFTIN1b 115  AAAAAAAA-----S--Y--A-----R--H----- 185
osRAFTIN1 154  NDDEPFGYD-----GSGTAASTT--V-TGA-----E--Y--A--TS-----R-----I--A-- 233

taRAFTIN1a 213  GVLATFGVADSATVASMEATLRACESPTIAGESKFCATSLEALVERAMEVLGTRDIRPVTSTLPRAGAPLQTYTVRSVR 292
taRAFTIN1b 186  I-----I-----T--P-----G-----VA-Q 265
osRAFTIN1 234  A--L-----P-T-EA-G-RE--T--W--L-----G--AA-----AALA-----G--A-A--A-L 313

taRAFTIN1a 293  PVEGGPFVEVACHDEAYPYTVYRCHTTGPSRAYMVMEGAAAARGGDAVTIATVCHTDTSLWNPEHVSFKLLGTPGGTPV 369
taRAFTIN1b 266  -----T-----AAA--A-----A----- 342
osRAFTIN1 314  ---AG--Q-----A-----E--DGGGD--E--V-----N--R-----S-- 392

taRAFTIN1a 370  CHLMPYGHIIWAKNVNRSPA 389
taRAFTIN1b 343  -----K----- 362
osRAFTIN1 393  -----V-----KS-T- 412

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B

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taRAFTIN1a 169  FFHEEAVRVGKRLPFRFPPTAAALGFLPRQVADSVPTTAALPGVLATFGVASDSATVASMEATLRACESPTIAGES 246
PG-bet 415  --R-KMLKS-TIMPM--DIKDKMPKRS--VI-SKL--S-SKIAELKAFHAGDE-QVEKMIGDA-SE--RAPS--T 492
RD22 176  --L-KDLVR--EMNV--NAEDGYGGKTA--GE-ET--GSEKFSET-KR-S-EAG-EEAEM-KK-IEE--ARKVSG-E 255
ASG-1 87  --N-HD-LE--TE-MY--SVTA--K--Q-R-VQEI--S-R-ADI--L-HIPPG-SEAADVAT--GL-DAAANGDVV 164
CFC1 124  --L-KDMHP-ATMSLH-TEN-AAAASA--Y-T-QKI--SSDK--EIFNK-S-KPG-LK-EM-KN-IKE--Q-A-E--E 200
SCB1 92  --L--DL-A--IFNMK-VNN--AA-TVPL--ISKQI--SEDKKKQ--ML--EAN-SNAKIIAE--GL-QE-ATEG-R 169

taRAFTIN1a 247  KFCATSLEALVERAMEVLGTRDIRPVTSTLPRAGAPLQTYTAAARSVRPVEGGPFVACHDEAYPYTVYRCHTTGPSRAY 324
PG-bet 493  -R-VN-A-DMIDFATS--RNVAA--RT-EDTKGSNGNIMIGSVKINGGKVTKSA--S--QTL--LL-Y--SVPKV-V- 569
RD22 256  -Y-----SM-DFSYSK--KYHV-A-STEVAKON--M-K-KIAAAG-KCLSDDKA--V--KQK--FA-FY--KAMMTTV- 334
ASG-1 165  RA-V--PDDM-G--AA--SNMQVLAPS--TG-MS--P--AA-A-K--D-SDA--G--P-L--S--SVQTGT- 241
CFC1 201  -Y-----SMIDYSISK--KV-AQA-STEVEKQATEM-K--IAAAG-QKMTDDKAA--V--KQN-A-A-FY--KSETT-- 276
SCB1 170  -H-----SM-DFVYSA--KAVGAFSTEKERETESAGKEV-VKNG--KLGDHVAIA--PMS--V-FG--LVPR-SG- 246

taRAFTIN1a 325  MVMDEGARGAGDAVTIATVCHTDTSLWNPEHVSFKLLGTPGGTPVCHLMPYGHIIW 380
PG-bet 570  EA-ILDPNSKVKINHGVAI--V--S-G-S-GA-VA--SG--KIE--WIFENDMTW 626
RD22 335  A-PL--AAEN-MRAKAVA--KN--A--N-LA--V-KV--TV--FL-ET-VV- 389
ASG-1 242  VME-QSSY-N-G-LKLVA--RN-TS-D-----V-AS--L-I--FV--V-F 298
CFC1 277  --PL--AAAD-TKAKAVA-----A--K-LA-QV-KVE--TI--FL-RD--V- 331
SCB1 247  L-RLK-AAED-VR-KAVVA--R--K-DHN-GA--V-NL--NGT--VFTE-NLL- 301

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Top
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Figure 4 Aiming Wang *et al.* 2002

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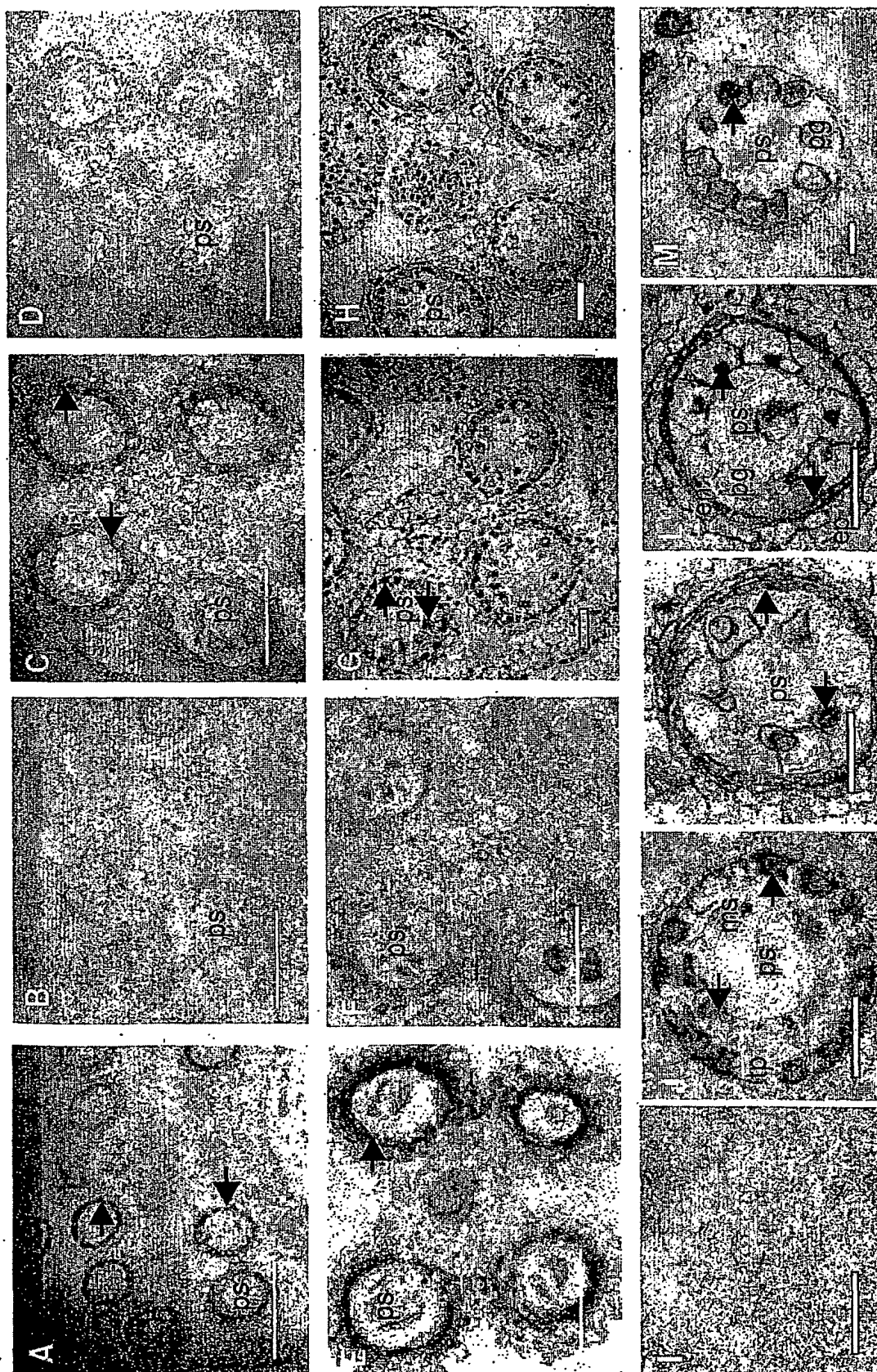


Figure 5

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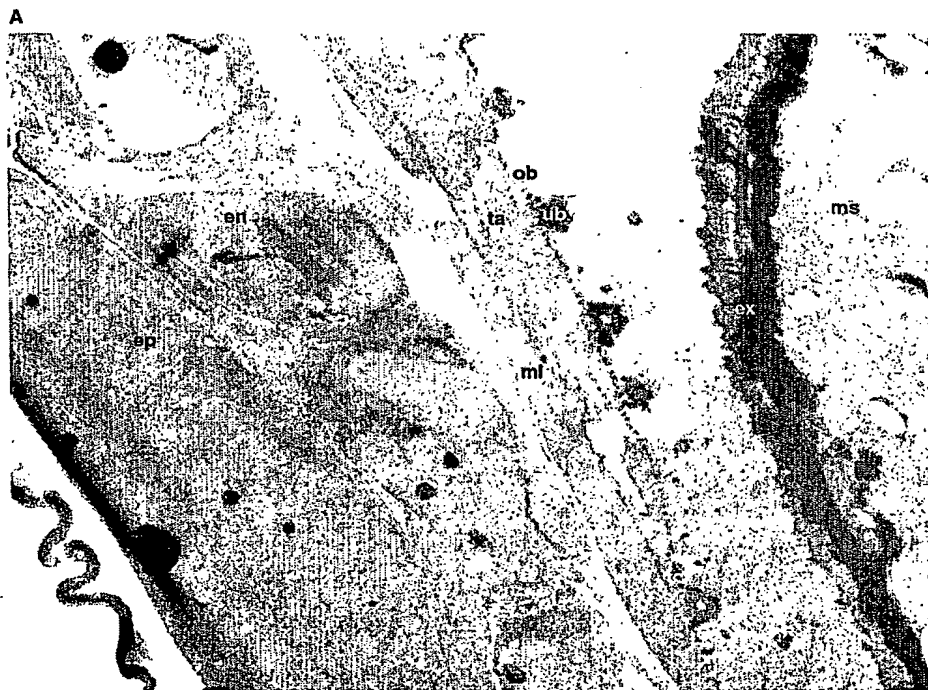
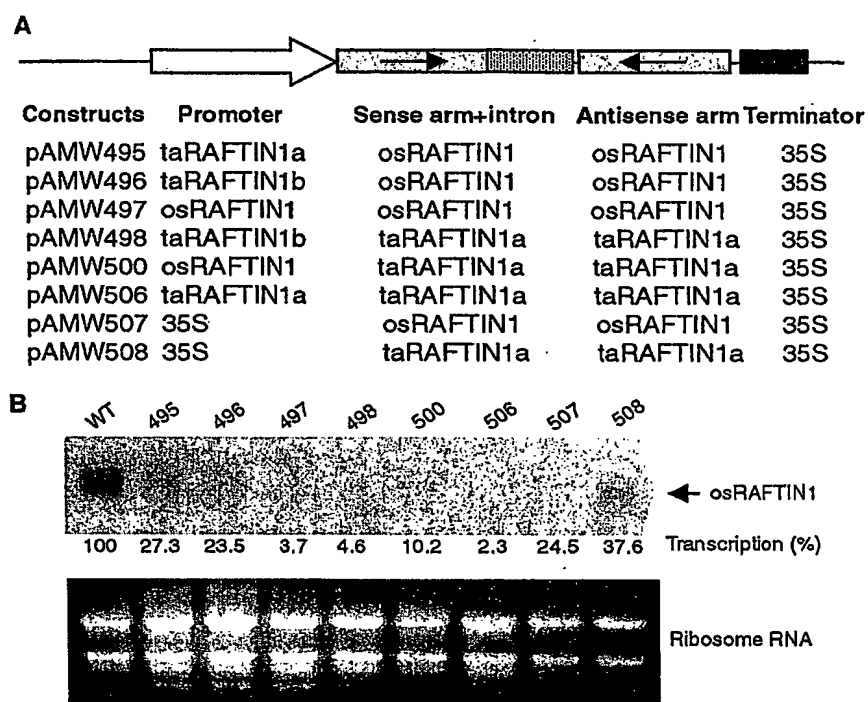


Figure 6

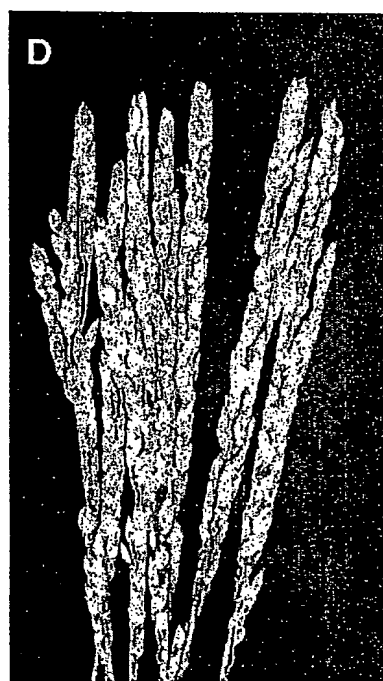
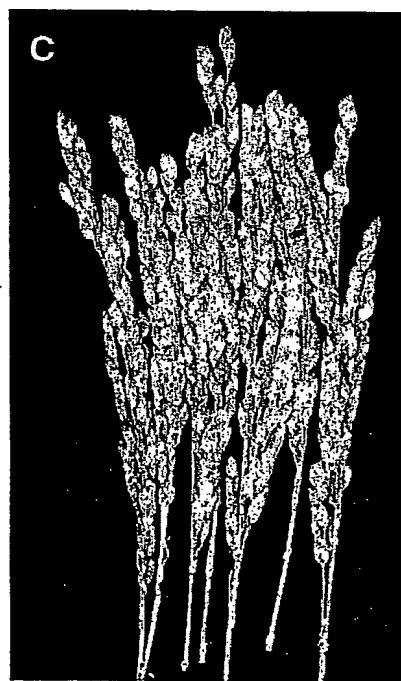
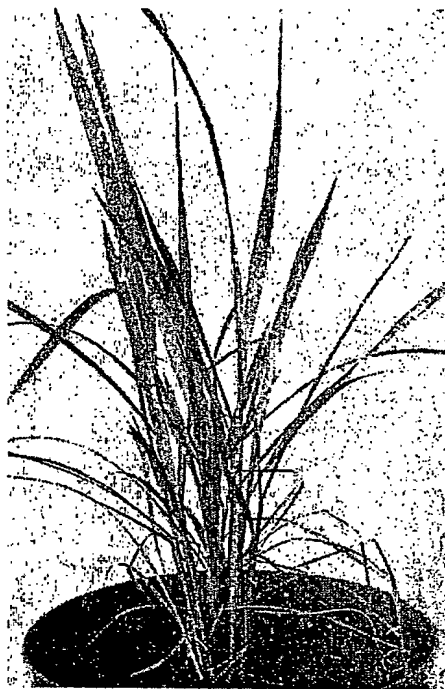
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Figure 7 Aiming Wang *et al.* 2002

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Figure 8ABCD Aiming Wang *et al.* 2002

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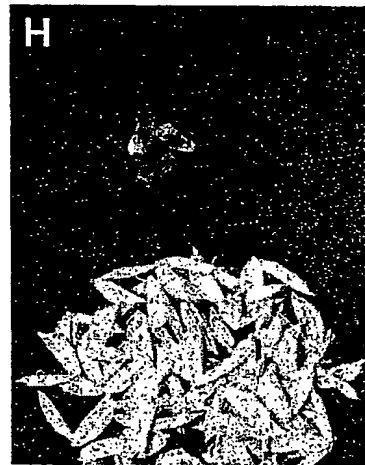
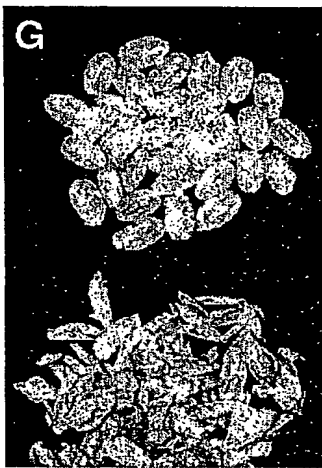
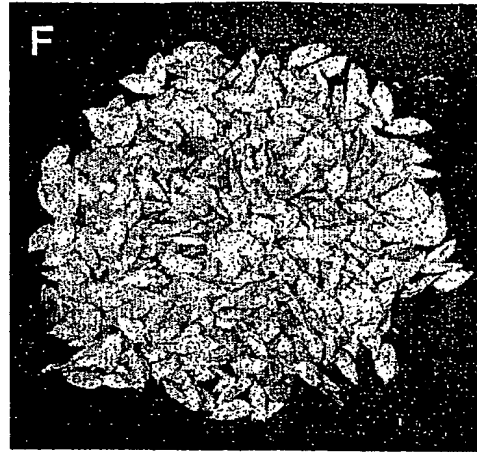
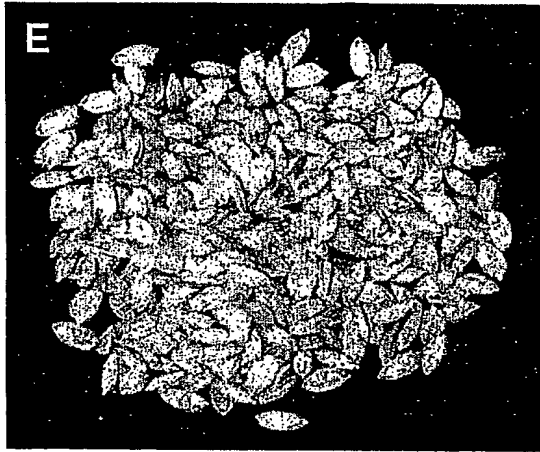


Figure 8EFGH Aiming Wang *et al.* 2002

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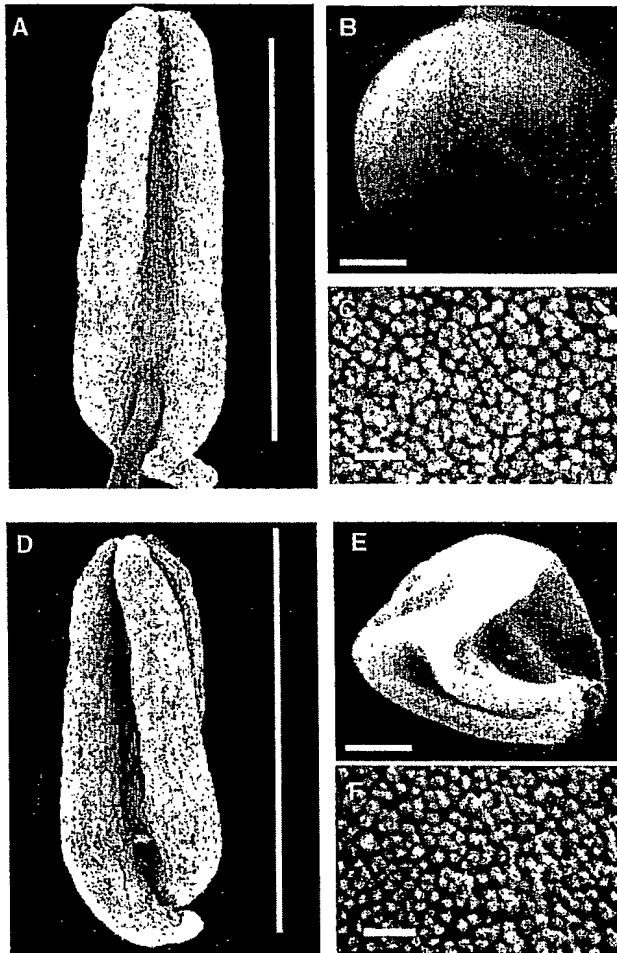
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Figure 8I Aiming Wang *et al.* 2002

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Figure 9 Aiming Wang *et al.* 2002

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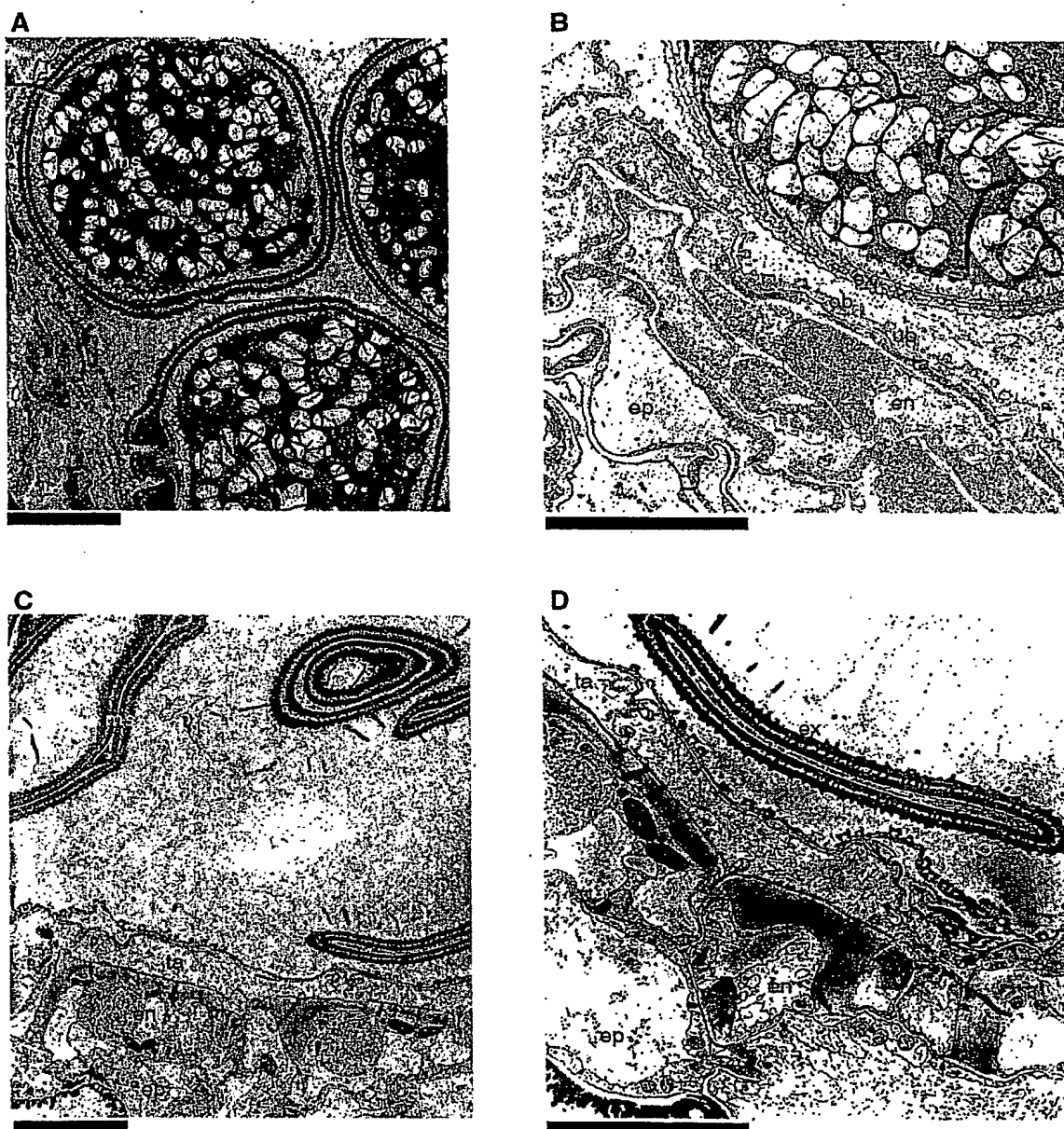


Figure 10

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Fig. 11. *taRAFTIN1a* cDNA sequence (1338 nt excluding the polyA tail, ORF from nt 29 to nt 1198). Start codon and stop codon are underlined.

CTCTGGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCCTCGCCACCAC
CCTGGTCGCGGTTCAAGGCTGGAGGGCAGCTGGGGCCACGCGGCGCCGGCGACGGCGGAGGTGTT
CTGGCGCGCCGTGCTGCCACACTCGCCATTGCCCGACGCCGTTCTCCGCCTTCTCAAACAACC
CGCAGCAGGTGTTGAACTGCTCACAGAAGCCACCAGCTTCGTGAGGGATGCCGAGGACAGGGCC
CCCCTTCGACTACCGTGATTACAGCCGCTCGCCGCCCGATGATGAACCGAGCAAGAGCACCGG
CGCCGCTCCGGGGCGCGGGACTTCGACTACGACGACTACAGCGGGGGCGACAAGCTCCGTGG
CGCCGCTCCGGGGCGCGGGACTTCGACTACGACGACTACAGCGGGGCCGACAAGCTCCGTGG
CGCCACCGATGAATACAAGGCGCCGAGCAGCAGCCTCGCTGGAAACGGGGCGTCCATGGCTAG
GGGCGGCAAGGCGGAGACGACGACGGTGTCTTTACGAGGAGGCGGTGCGCGTCGGCAAGAG
GCTCCCATTTCCGCTTCCCGCCGGCGACTCCCGCCGCGCTCGGTTTCTGCGCGCCAGGTGCG
CGACTCCGTCCCGTTACGACGGCCGCGCTGCCTGGCGTCCTCGCGACGTTGGCGTCGCGTC
CGACTCCGCCACGGTGGCCAGCATGGAGGCGACGCTGCGCGCCTGCGAGTCGCCGACCATCGC
CGGGGAGTCCAAGTTCTGCGCGACCTCGCTGGAGGCCCTGGTGGAGCGCGCCATGGAAGTGCT
GGGGACCCGCGACATCAGGCCGGTGACGTCGACGCTGCCCCGCGCCGGCGCCCCGCTGCAGAC
GTACACCGTCCGCTCCGTGCGGCCGGTGGAGGGGGGCCCTGTCTTCGTGGCGTGCCACGACGA
GGCCTACCCGTACACCGTGTACCGGTGCCACACCACTGGCCCGTCCAGGGCGTACATGGTGGA
CATGGAGGGCGCGCGCGGGCGGCGACGCGGTGACCATCGCCACCGTGTGCCACACCGACACGTC
CCTGTGGAACCCGGAGCACGTCTCCTTCAAGCTCCTGGGCACCAAGCCTGGCGGCACGCCGGT
CTGCCACCTCATGCCGTACGGGCACATAATCTGGGCCAAGAACGTGAATCGCTCGCCGGCGTG
AGCGGGCCCGGGCAGCTCTGTGGTCTCGCCGGAACTAAGATCGATGTACTACTACTATCTG
TTTCCACCTACGTCTTCTGTTGTTTCAGACCACCAGATGGTCACCAGAGCAGCGCTTGTAATAA
AAGAACAGCTTCTGCAAAAAAAAAAAAAAAAAA

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Fig. 12. *taRAFTIN1a* genomic sequence (1560 bps including two introns). Introns are shown in lower case letters. Start codon and stop codon are underlined.

CTCTGGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCCTCGCCACCAC
CCTGGTCGCGgtaatggccgaagaagccactgagcaacgcctgcatcttcttcatttcggcaa
actgcacctagtgcatttcgcatgagattgatcgatcacaaactggtgctaacggcctgtttc
gtcacagGTTCAGGCTGGAGGGCAGCTGGGCCACGCGGCGCCGGCGACGGCGGAGGTGTTCTG
GCGCGCCGTGCTGCCACACTCGCCATTGCCCGACGCCGTTCTCCGCCTTCTCAAACAACCCGC
AGCAGgtctgtctttcatgttcccttccctcgtcgcccctecgttaactgtcttcttctctcgag
tttgattgaccgccaaacacaaaaaatgcatgcacgcacagGTGTTGAACTGCTCACAGAAGC
CACCAGCTTCGTGAGGGATGCCGAGGACAGGCCCCCTTCGACTACCGTGATTACAGCCGCTC
GCCGCCCGATGATGAACCGAGCAAGAGCACCGGCGCCGCCTCCGGGGCGCGGGACTTCGACTA
CGACGACTACAGCGGGGGCGACAAGCTCCGTGGCGCCACCGATGAATACAAGGCGCCGAGCAG
CAGCCTCGCTGGAAACGGGGCGTCCATGGCTAGGGGCGGCAAGGCGGAGACGACGACGGTGTT
CTTTCACGAGGAGGCGGTGCGCGTCGGCAAGAGGCTCCCATTCCGCTTCCCGCCGGCGACTCC
CGCCGCGCTCGGTTTCTTGCCGCGCCAGGTGCGCCGACTCCGTCCCGTTCACGACGGCCGCGCT
GCCTGGCGTCTCTCGCGACGTTGCGCGTCGCGTCCGACTCCGCCACGGTGGCCAGCATGGAGGC
GACGCTGCGCGCCTGCGAGTCGCCGACCATCGCCGGGGAGTCCAAGTTCTGCGCGACCTCGCT
GGAGGCCCTGGTGGAGCGCGCCATGGAAGTGCTGGGGACCCGCGACATCAGGCCGGTGACGTC
GACGCTGCCCCGCGCCGGCGCCCCGCTGCAGACGTACACCGTCCGCTCCGTGCGGCCGGTGGA
GGGGGGGCTGTCTTCGTGGCGTGCCACGACGAGGCCTACCCGTACACCGTGTACCGGTGCCA
CACCCTGGCCCCGTCCAGGGCGTACATGGTGGACATGGAGGGCGCGCGCGGCGGCGACGCGGT
GACCATCGCCACCGTGTGCCACACCGACACGTCCCTGTGGAACCCGGAGCACGTCTCCTTCAA
GCTCCTGGGCACCAAGCCTGGCGGCACGCCGGTCTGCCACCTCATGCCGTACGGGCACATAAT
CTGGGCCAAGAACGTGAATCGCTCGCCGGCGTGAGCGGCCCGGGCAGCTCTGTGGTCTCGCCG
GAACTAAGATCGATGTACTACTACTATCTGTTTCCACCTACGTCTTCTGTTGTTTCAGACC
ACCAGATGGTCACCAGAGCAGCGCTTGTAATAAAAGAACAGCTTCTGC

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Fig.13. *taRAFTIN1a* promoter sequence (1719 bps).

CTGTGCGATGGCGCTCTGTCCTTGTGATTCTTTCTTAGGGAACTCGTCTCTGGGGCCTCCGAGG
CCTGCAACCCTGTATCAGGACAATTCTGACTGGCCTCCAGGAGTCCTAACAGCCACCGACCTG
GTCCACTGGGCCCATCTAGAGTATCTTGAAGTGTCGTTTGACAAATCCCGCTAATTAAGGGA
TGTGATGATGATGGTTTCTGAATCCGCGCGCCTTACCTCGCAAAACGGGGAATTGCAAAGGAT
ATATGGCACCTGTCGCGTCGTGAGGCCAGACGCTTCGGTTTCAAGCTGGTTATAGGGAGGGGG
AAACGAAGGGTTTTTTCTCCCTCTGTCTTCATCCATTTTCGTCTCCCAGCCCTCAGCTCCCAA
AAGCGTGTCGCCACCTCAAAGTCTTCAGCGCTTGCTCACGTAGCCCCCGTCCACCCCTTCCTT
GCCACCAAGATGGCCCGAACCAAGAGCGAGAAGGTTCCCTAAGGTTCCCAGCTAGGATCTGCCC
GCCGCTGGAACGGGGCTGAAGCGGAAGAGGGTGCCTCCAAGGGTGGTATGAAACAACAGCCG
GAAGCCCCCAAGACTACAGGAAAGTGGTTCCTTCCTCGGCCACCGACAAAAAACTTCAGGGT
CTCGTGGAGATAGGGCTGATGCCAGCGGATTTGGAGTGCCGCCTCCCGGGGGACGAGGCTCCG
CCAACCTCCTCGCGACGGTGAGCACATCCTCTGCCTGGAGTATATATTTTCGGGAGGGGGCTCGGG
TTTCCCTACACGACTTCGTTTGCGGGATCTTGCGCTTCTACGGCTGCTAGCTACACCACATC
CCGTCAAACGGGGTTCTTTACATTGCAAACCTTCATCACATTTTGCGAGTGCTTTCTCGGGACT
GCCGCTCACTTTAAGTTGTTCCAATACTTCAATCAGGACTGCGTTCAGACCAACGGGGACATC
GTCTACGACCCCGCAACACCAAATTCCTCGCCACATACCTCCGGAAAATAATCCTATACAACC
TGGTCTCACGCTTCATCTCGTAAGATTTGCCATGTGTACTTCACCAATCTTGATGCATCCCTT
TTTCCCAAGATTTATATGCCTGATCTGTATTTTGTCTCCGCTGTTTCGAGATTTGATGTTTA
ATTGATGAAGCCCAAGCAATCCGGCATGCCCGTCGGTGCACTAGATGGCTAGCTTTTCTACGG
TGCTGGGCCTGCCGCGAGGGGCGCGAGGCCACGTAGGAGACTGTTAGGATTCATGGGGCTGG
ACGCTGGTGGCGTGAAGTTCGGGAAGGAGGATTGAGGAAGAAGGATGCATCAAGATTGGTGAA
GAACACGTGGCATCCTCTAGAGTAGGTCTTACGAGATGAAGCCTGAGACCAGGTCGTATGGGA
TTATTTTCCCGACCTCCCGAAGCCCCGCAAAGTTAACTGCAGCTGCGTGGACGGCGAGCACC
GCACCGCACACGAACGCGAACCTGACGCTGCCGCGCCACACAACACGCCATTTCGCGCGCGGAT
CGTCGGATGTCACGCCCAGGATTATATTCTCCGGTGCCGCACGTACCATGCGATCGCACAGCT
CACGTCGAGAGCTTTTCTGTTTGCGGTCGCCGTCAATGAAACACCTTCCCGTCGAGCCGACGA
CGCCTATAAGTACCTCGTCTGATCGCATCATCACTCCCAAGTACTACAACCTCTGGACCTCTC
ACCTAGCGCACATCCATG

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Fig.14. *taRAFTIN1b* cDNA sequence (1275 bps excluding the polyA tail, ORF from nt 25 to nt 1113). Start codon and stop codon are underlined.

CGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCCTCGCTGCCACCCTG
GTCGCGGTTTCAGGCTGGAGGGCAGCTGGGCCACGCGGCGCCGGCTACGGGGGAGGTGTTCTGG
CGCGCCGTGCTGCCGCACTCGCCATTGCCTGACGCCGTTCTCCGCCCTCCTCAAACAACCTGCA
GCAGAATCCACCAGCTTCGTGAGAGACCCCGAGGACAGGCCCCCTTCGACTACCGTGATTAC
AGCCGCTCGTCGTCCGATGATGAACCGAGCAAGAGCACCGTTCGCCGCCTCCGGAGCGGGGGC
TTCGACTACGACAACTACAGCGGGGCGGACGAACGTCGTGGTGCCACCGATGAATACAAGGCG
CCGAGCAGCAGCCTCGCTGGAAGCGGGGCGTACATGGCTAGGGGCGGCAAGGCGGAGACGACG
ACGGTGTTCTTTACGAGGAGGCGGTGCGCGTCGGCAGGAGGCTCCCATTCACCTTCCCGCCG
GCGACTCCCGCCGCTCTCGGTTTCTTGCCGCGCCAGGTGCGCGACTCCGTCCCGTTCACGACG
GCCGCGCTGCCCCGGCATCCTCGCGACGTTTGCGCATCGCGTCCGACTCCACCACGGTGCCCAGC
ATGGAGGCGACGCTGCGCGCCTGCGAGTCGCCCACCATCGCCGGGGAGTCCAAGTTCTGCGCG
ACTTCGCTGGAGGCCCTGGTGGAGCGCGCCATGGGAGTGCTGGGGACCCGGGACATCAGGCCG
GTGACGTCGACGCTGCCCCGCGCCGGCGCCCCGCTGCAGACGTACACCGTCGTGCGCCGTGCAG
CCGGTGGAGGGGGGGCCTGTCTTCGTGGCGTGCCACGACGAGGCCTACCCGTACACCGTGTAC
CGGTGCCACACCACCGGCCCGTCCAGGGCGTACACGGTGGACATGGAGGGCGCGCGCGGCCG
GACGCGGTGACCATCGCCGCCGTGTGCCACACCGACACGTCCCTGTGGAACCCGGAGCACGTC
TCCTTCAAGCTCCTCGGCACCAAGCCCGGCGGCACGCCGGTCTGCCACCTCATGCCGTACGGG
CACATAATCTGGGCCAAGAACGTGAAGCGCTCGCCGGCGTGAGCGGCCTTGACGCTCTGTGGT
GTCGCCGGAATAAGATCGATGTACTACTATCTGTTCTTACCTACGTCTTCTTGTGTTTC
ATACCACCAGATGGTCAACCAAGAGCAAGCGTTCGTAATAAAAAGAACAGCTTTTTTGCAGAAG
CTGGTGTTTTATTTTAAAAA

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Fig.15. *taRAFTIN1b* genomic sequence (1503 bps including two introns). Introns are shown in lower case letters. Start codon and stop codon are underlined.

CGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCCTCGCTGCCACCCTG
GTCGCGgtaatggccgaagaagagcaacgcctgcatcttcttcattttggcaaatgacaccta
gtacattttgcatgagattaatcaatcacaaactggtgctaacggcctgtttcgtcccag
G TTCAGGCTGGAGGGCAGCTGGGCCACGCGCGCCGGCTACGGGGGAGGTGTTCTGGCGCGCC
GTGCTGCCGCACTCGCCATTGCCTGACGCCGTTCTCCGCCCTCCTCAAACAACCTGCAGCAGgt
ctgtcttgcattcttctcgtcgccctccgttaactgtcttcttctctcgagtttgattgatca
ccaaacacaaaaatgcatgcacgcgtgggtgttgaaactgcgcacagAATCCACCAGCTTCGTG
AGAGACCCCGAGGACAGGCCCCCTTCGACTACCGTGATTACAGCCGCTCGTCGTCCGATGAT
GAACCGAGCAAGAGCACCGTCGCCGCCTCCGGAGCGGGGGGCTTCGACTACGACAACTACAGC
GGGGCCGACGAACGTCGTGGTGCCACCGATGAATACAAGGCGCCGAGCAGCAGCCTCGCTGGA
AGCGGGGCGTACATGGCTAGGGGCGGCAAGGCGGAGACGACGACGGTGTTCTTTACGAGGAG
GCGGTGCGCGTCGGCAGGAGGCTCCCATTCCACTTCCCGCCGCGACTCCCGCCGCTCTCGGT
TTCCTGCCGCGCCAGGTGCGCGACTCCGTCCCGTTACGACGGCCGCGCTGCCCGGCATCCTC
GCGACGTTTGGCATCGCGTCCGACTCCACCACGGTGCCAGCATGGAGGCGACGCTGCGCGCC
TGCGAGTCGCCACCATCGCCGGGGAGTCCAAGTTCTGCGCGACTTCGCTGGAGGCCCTGGTG
GAGCGCGCCATGGGAGTGCTGGGGACCCGGGACATCAGGCCGGTGACGTGACGCTGCCCCGC
GCCGGCGCCCCGCTGCAGACGTACACCGTCGTGCGCGTGACGCCGGTGGAGGGGGGGCCTGTC
TTCGTGGCGTGCCACGACGAGGCCTACCCGTACACCGTGTACCGGTGCCACACCACCGGCCCG
TCCAGGGCGTACACGGTGGACATGGAGGGCGCGCGCGCGCGGACGCGGTGACCATCGCCGCC
GTGTGCCACACCGACACGTCCCTGTGGAACCCGGAGCACGTCTCCTTCAAGCTCCTCGGCACC
AAGCCCGGCGGCACGCCGGTCTGCCACCTCATGCCGTACGGGCACATAATCTGGGCCAAGAAC
GTGAAGCGCTCGCCGGCGTGAGCGGCCTTGACGCTCTGTGGTGTGCGCCGGAATAAGATCGAT
GTACTACTACTATCTGTTCTTACCTACGTCTTCTTGTGTTTCATACCACCAGATGGTCACCCA
AGAGCAAGCGTTCGTAATAAAAAGAACAGCTTTTTGCAGAAGCTGGTGTTTTATTTT

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Fig. 16. *taRAFTIN1b* promoter sequence (2095 bps).

TTGTTGAGTGCCACACTATATTCACTACACCATATGCACATTATGCTTGGATTGTCTTGTACT
TGACTCATGTGTTTAGACACTTCATTTTATTTGGTGTGTGAATGACTCCTATGCTTACCATA
GACCTTTTCATTGAGCGCTTTGTGCATGTTTGTATACCTTGAGGTAGATGTTTGTCTCTTGT
CAAATATATAGCATCTCTACCTCCCATTTCATGCTTGTTCATGATGTCCTTGATTGTGCT
CAATTCATATGCTTCTGTGACATGCCACAATCCTTTGTACACCATATGCTAGGCTTGATGAT
GACACTTGTGTTGGGTGACTCACCTTTTGAATGATTGGTTCATTAACGCTAACCACATTTAT
TTTTCCAAGTGTTTGTTCCTTGCTCCTTTTGAAGGAACCACATGACGGTTCGACATTGGAG
AGTGCCTATTTTCGAGCTTCAAGATGATGAGTGTGCTTGGTGATCGTCCACTTCTACATGGTGACG
CCGTCTCTTTCCCATGGTGATTGGTTCCTGATCCGAGGTTCGGATCTTTCCCAAGTGGGAGGG
GATGATGCGGAGCATACTACGGACATCACCATGTCTAGAGTTCATTTCAGCAAGTGACACCTAT
CACATCTACTTCACATACATAAAGGTGAATCATCTCCTTTACACGTGCTCACTTGATCCCTTC
GAGGATGGTATACTACTTGACACTTCTCACGTGTGCATGCATAGGCATTGTTCGGAGCACCATG
AACGATGAGGAGGAGTGCGAGCACAAGTGTACAACCTACACCATCCGCGAGGGAAGCATGGAAG
AGAAGGAAGAAGAAGCATGGACAAGCTTCTGGAAAGCCCGGAACCTTCTGGCCTCCTGCCCGGA
ACTTCCGGTCATCCGAAACTTCCTGCCCGGACACACCGAAGCCGTCTGAGAGCGTGCCAAATC
TCTGGATAGCCCGGACCTTCGACCGGAACCTCCGGCGCCTGGACCTTCCGGCCATCCCTGGAA
CTCCCGCCTGCCTGCACGCAGAGACTCGGGCCGAAGCGCATGTACCCTTTCGCCCTCACTT
ATCCCTTCGTGGCTATCACTATATATACTCATCCTCCTCCTCCATTCTAGGGTTAGCATTTTG
ATAGCTCATTTGCATGTGAGATTTGCTCCTTACCCCCATCTCCTCTTGAGAGAGTGAGATTGA
TGCACTCCATTGGAGTCCAAGGTCTCCTTTGGAGAAGATCCCATAGGGGAATCAAGACCCCAT
CATGGGAAGATCCTTCTAGGATTCAAGACCTCAACTCCTTTAAGGATTGGGATGAACTAGTTA
CCTCTTGTATCTTCTTGTGTTGGATTTAAACCTTTGTATCCCTCTATGTGTATGTGGATTTAG
CATATGTGTGATTGGATCTTGTCTATTGGAGTGTTCCTCTCTTTTGTTCCTTGTGTTCAT
CGTTTTCTTCGGGAGATCCCCTCCATTTCTGTAAAGATCGGTCCCTAGGGTTCTACCCTACAT
TAGCTCAGGTTTCCCCTACACATCTTCGTTTGTGAGCTGTTGCGCTTCTACGGCTGGGAGCTA
CAGCACATCTCATTCCCACCAAACGGGGTTCTTCACATTGTAAACTTCATCGTATTTTGCGAA
TGCTTTCTGGGGACAGCCACTCACTTTGAGTTGTTCCGATACTTCTTCCGGGTCTGCGTTTCAG
ACCAACGGGGACACCGTCTGCAACCTTGGAGGAGCCATTCTTCCGACACACCAAAATTTTCGC
CACGGACCCCCCGAAGATCCGCAAGAAAAAAGCTGCAACGGCGTGGACGGCGAGCACC GC
ACCGCACACGAACGCGAACGCGACGCTGCCGCGCCACACAACACGCCATTTCGCGCGCGGATCG
TCGGATGTCACGCCACGATAATATTCTCCGGTGCCGACGTACCATGCGATCGCACAGCTCA
CATCGAGAGCTTTTCTGTTTGGTGTGCGCGTCAATGAAACACCTTCCCGTCAAGCCGACGACG
CCTATAAGTACCTCGCCTGATCGCATTATCACTCCCAAGTACTACAACCTCTCGACCTCTCAC
CTAGCGCACATCCATG

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Fig. 17. *taRAFTIN1d* predicted cDNA sequence (246 bps).

ATGGCGCGCTTCCTCGTCGCCCTCCTCGCTGCCACCCTGGTCGCGGTTTCAGGCTGGAGGGCAG
CTGGGCCACGCAGCGCCGGCGACGGCGGAGGTGTTCTGGCGCGCCGTGCTGCCGCACTCGCCA
TTGCCCCGACGCCGTTCTCCGCCTCCTCAAACAACCTGCAGCAGGTGTTGAACTGCACACAGAA
GCCACCAGCTTCGTAAGAGACCCCGAGGACAGGCCCCCCTTCGACTACCGTGATTAC

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Fig. 18. *taRAFTIN1d* partial genomic sequence (441 bps). Introns are shown in lower case letters.

ATGGCGCGCTTCCTCGTCGCCCTCCTCGCTGCCACCCTGGTCGCGgtaatggccgaagaagcc
actgagcaacgcctgcatcttctttattttggcaaactgggtgctaacggccaatactgccgct
tgcgttacgtctcagGTTCAAGGCTGGAGGGCAGCTGGGCCACGCAGCGCCGGCGACGGCGGAG
GTGTTCTGGCGCGCCGTGCTGCCGCACTCGCCATTGCCCGACGCCGTTCTCCGCCTCCTCAAA
CAACCTGCAGCAGgtctgtcttgcatgttcctcgtcgccctccgttaactgtcttcttctctc
gagtttgattgatcaccaaacacaaaaatgcatgcacgcgtacgcgtagGTGTTGAACTGCAC
ACAGAAGCCACCAGCTTCGTAAGAGACCCCGAGGACAGGCCCCCTTCGACTACCGTGATTAC

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Fig. 19. *osRAFTINI* cDNA (1301 bps, ORF from nt 63 to nt1301). Start and stop codons are underlined.

GTCGCAGTCGTCTCCGGCGAGAAATCGGCTGCGCCCCGTCTCTCTCTCTCTCGAACGCTTCCA
TGGCGCGCTTCCTCCTCCTCCTCGTCGCCGCTGCCGCGCCGCTGCTTTCGCTGGGCG
ACGCGGCGCCGTCGACGGCCGAGGTGTTCTGGCGCGCCGTGCTGCCGGAATCCCCGTTGCCGG
ACGCCTTCCTCCGCCTCCTCCGCCCTGACACCAGCTTCGTCTGTCGGCAAAGCGGAGGCGGCCG
GTGGCGCGGCGCGGACCGGATTCCCCTTCGATTACACTGACTACAGGGGATCTGATTCTCCGA
CGACGGCGAGTGGTTTGGACCTCGCCGGTGACTTCGGCGAGCCGGCGCCTTTTCGGCTACGACT
ACAGTGCACAGGGCGAAGGCGGCGGCGGCGGCCGCCGCCGCCGCCGGGAGAGCAGGTTCTTG
CCGTTCGACGCGGGCTTCAACTACGACAAATACGTCCGCGCGAGGAAGCTCCGCGGCGGCAGCA
GCACCGCCGGCGGAGAGAATGATGACGAGCCTTTCGGGTACGACTACAAGGCGCCGAGCAGCG
GCAGCGGCACCGCGGCGTCGACGACGGCGCGAGGCGTCGGCACGGGCGCCACGACGACGGTGT
TCTTCCACGAGGAGGCGGTGCGCGTCGGCGAGAGGCTCCCGTTCTACTTCCCGGCGGCGACGA
CGTCGGCGCTGGGCTTCCTGCCGCGCCGCGTCGCGGACTCCATCCCGTTACGGCGGCGCGCGC
TGCCGGCCGTCCTCGCGCTGTTTCGGCGTCGCGCCGGACACCGCCGAGGCGGCGGCGCATGAGGG
AGACGCTGCGCACGTGCGAGTGGCCGACCCTCGCCGGCGAGTCCAAGTTCTGCGCCACGTGCG
TGGAGGCCCTGGTGGAGGGCGCCATGGCGGCGCTCGGGACACGCGACATCGCCGCGCTGGCGT
CGACGCTGCCCCGCGGCGGCGCGCCGCTGCAGGCGTACGCCGTCCGCGCCGTGCTCCCCGTCG
AGGGCGCCGGCTTCGTGGCGTGCCACGACCAGGCGTACCCGTACACCGTGTACCGCTGCCACA
CCACCGGCCCGGCCAGAGCTTACATGGTGGAGATGGAAGGCGACGGCGGCGGCGATGGCGGCG
AGGCGGTGACCGTGGCCACCGTGTGCCACACCAACACGTGCGCGGTGGAACCCGGAGCACGTCT
CGTTCAAGCTCCTCGGCACCAAGCCCGGCGGCTCGCCGGTGTGCCACCTCATGCCGTACGGGC
ACATCGTCTGGGCCAAGAACGTGAAGAGCTCGACGGCGTAG

GTCGCAGTCTCTCCGGCGAGAAATCGGCTGCGCCCCGTCTCTCTCTCTCTCGAACGCTTCCA
TGGCGCGCTTCTCTCTCTCTCTCGTCGCCGCTGCCGCCGCCGTGCTTTTCGgtacact
catgatgccgctactcagctgagccatgcaccggtgcacccgtataactaacgatcgctcgatc
gaccgacgatgtgtgtttcttcagcagCTGGGCGACGCGGCCCGTCGACGGCCGAGGTGTTCT
GGCGCGCCGTGCTGCCGGAATCCCCGTTGCCGGACGCCCTTCTCCGCCTCTCCGCCCTGgtc
ggtgtccttctcttctctctcttcgcccgcgcgcgcgcgcgcattactctcctcgagggtttgatttg
tttggtggacggttgcagACACCAGCTTTCGTCGTCGGCAAAGCGGAGGCGGCCGGTGGCGCGGCG
CGGACCGGATTCCTTTCGATTACACTGACTACAGGGGATCTGATTCTCCGACGACGGCGAGT
GGTTTGACCTCGCCGGTGACTTCGGCGAGCCGGCGCCCTTTCGGCTACGACTACAGTGCACAG
GGCGAAGGCGGCGGCGGCGGCGCCGCCGCCGCCGCCGGGAGAGCAGGTTCTTGCCGTCGACGCG
GGCTTCAACTACGACAAATACGTCGGCGCGAGGAAGCTCCGCGGCGGCAGCAGCACCGCCGGC
GGAGAGAATGATGACGAGCCTTTCGGGTACGACTACAAGGCGCCGAGCAGCGGCAGCGGCACC
GCGGCGTCGACGACGGCGCGAGGCGTCGGCACGGGCGCCACGACGACGGTGTTCTTCCACGAG
GAGGCGGTGCGCGTCGGCGAGAGGCTCCCGTTCTACTTCCC GGCGGCGACGACGTCGGCGCTG
GGCTTCTTGCCGCGCCGCGTCGCGGACTCCATCCCGTTACGGCGGCGCGCTGCCGGCCGTC
CTCGCGCTGTTTCGGCGTCGCGCCGGACACCGCCGAGGCGGCCGGCATGAGGGAGACGCTGCGC
ACGTGCGAGTGGCCGACCCCTCGCCGGCGAGTCCAAGTTCTGCGCCACGTCGCTGGAGGCCCTG
GTGGAGGGCGCCATGGCGGCGCTCGGGACACGCGACATCGCCGCGCTGGCGTCGACGCTGCCC
CGCGGCGGCGCGCGCTGCAGGCGTACCGCTCCGCGCCGCTGCTCCCCGTCGAGGGCGCCGGC
TTCGTGGCGTGCCACGACCAGGCGTACCCGTACACCGTGTAACGCTGCCACACACCAGCGCCG
GCCAGAGCTTACATGGTGGAGATGGAAGGCGACGGCGGCGCGCATGGCGGCGAGGCGGTGACC
GTGGCCACCGTGTCGACACCAACACGTGCGCGTGGAACCCGGAGCACGTCTCGTTCAAGCTC
CTCGGCACCAAGCCCGGCGGCTCGCCGGTGTCGACCTCATGCCGTACGGGCACATCGTCTGG
GCCAAGAACGTGAAGAGCTCGACGGCGTAG

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Fig. 21. *osRAFTIN1* promoter sequence (1461 bps).

CGAAGGCCAACTCTGGTAAGGATTCCCATTTACACGAATCAATTTAATAAGTCTAAAACGAACA
CTATGTTATGAGAAACACCTCACATCCGTCCATAACCGTGCGCATGACTATTTAAAAAGTTTA
ACTAAACTCTACAAAAGTTGCACGCTTTACCCACACGTCATGAACGTTTCACATTACCGAATA
CATGTGGATCGGACATGGCCGACAAAGGAGAGTTCAATACAAGGCTTTTCCATAACCAATCCA
TAAATATCCTATGTCCACGCTTGGGTGGAATCTCTCCACCAAACATCAAGCCAGGATCAGGT
CCTCATCTACCCATGCCCCACTCCATGGACTCCGACACATCCCCACTGCAGGAGATTGCCATA
TACGCCACCATAACAGTGCTCCTCAACCGCTAACATGTTGGACACCAAATTCTATATACTTAT
ATAGTTCATCTCCACTAAGTGTAAGTTAATTACATTTCTCTCTCTCTCATTAAGCCACATCAC
CTCAATTATTTTATAGCCTTTAGATGATAGATCTATGGTCCAAATTGTCTTTTCTTTCTTCTCT
CTTAAAAACATGCAATCTTAAATACTTTTAGGCTCAAAATTGTATCAAATTGTTTTAGTTTTG
TACATATTATGCAACTTAATTTTTCGCCGCAACGCGGAGGGGTATTTTCATCTAGTATTATTTA
AGAGCTATACACACTGCTATAGGGGAAAAAAGATAGGTTTGGCCCCCTGGTCAGTCTGT
GCACGGCTATATGTTGAAGGAAAAAGCCAGTACGTTTTGTAGGTTGTTTTTTTTTTAGAAAT
GCTAAAAAGTTGTGGCATGTTTTTTAGGTAAAAGCCTTTAAATATAAGTTACATTGTAAGTAC
AGTGTAATTCGCTGTAAGTATATTTGTAATCTCTATATAAGTTAGATATAAAATTACATATAT
ATTATTTTAATACTTATTTATAAGTTAGTATATTATAGTTATAATGGAATTAATTATAATTAT
AGTATAGTTAGATTTGAAAGTTTTTCCTTTAAGAAATTTTCGCAACAGTTTATTAGATATAGTC
CCTAAACGAAAATGTCAGGTGGATGCATGATTTCAGTGTGACGCTCGGGCGGATCACGGCTGCG
TCACGAAAATTCCTCCCATGCAACCCGCGTCCGGCCGTCCTTCGTGCCAACAGGCAACAGCGC
GGCGCCGGCGAACGTCACGCCCAAGATTATATTCCCCCTCTCGCGCTCGCGCGCGCCGCGACG
TCGTCCGAGCCAACATTATTTTTCTGTTTCTGTCAACGTCGCCGTTGATCTCAAGCGAGATT
TGAGGTTTGGCCACGACGACGCTGCCTATAAATACCAGGTGGTGGTCACCGCCCGGCGGCGT
CGATCGATCCGTCGCGAGTCGTCTCCGGCGAGAAATCGGCTGCGCCCCGTCTCTCTCTCTCG
AACGCTTCCATG

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Fig. 22. Predicted protein sequences

taRAFTIN1a (389 residues)

MARFLVALLATTILVAVQAGGQLGHAAPATAEVFWRAVLPHSPLPDAVLRLLLKQPAAGVELLTEATSFVR
DAEDRPPFDYRDYSRSPDDEPSKSTGAASGARDFDYDDYSGGDKLRGAASGARDFDYDDYSGADKLRG
ATDEYKAPSSSLAGNGASMARGGKAETTTVFFHEEAVRVGKRLPFRFPATPAALGFLPRQVADSVFPT
TAALPGVLATFGVASDSATVASMEATLRACESPTIAGESKFCATSLEALVERAMEVLGTRDIRPVTSTL
PRAGAPLQTYTVRSVRPVEGGPVFVACHDEAYPYTVYRCHTTGPSRAYMVDMEGARGGDAVTIATVCHT
DTSLWNPEHVSFKLLGTPGGTPVCHLMPYGHIIWAKNVNRSPA

taRAFTIN1b (362 residues)

MARFLVALLAATLVAVQAGGQLGHAAPATGEVFWRAVLPHSPLPDAVLRLLLKQPAAESTSFVRDPEDRP
PFDYRDYSRSSSDDEPSKSTVAASGAGGFDYDNYSGADERRGATDEYKAPSSSLAGSGAYMARGGKAET
TTVFFHEEAVRVGRRLPFHFPPATPAALGFLPRQVADSVFPTTAALPGILATFGIASDSTTVPSMEATL
RACESPTIAGESKFCATSLEALVERAMGVLGTRDIRPVTSTLPRAGAPLQTYTVAVVQPVVEGGPVFVAC
HDEAYPYTVYRCHTTGPSRAYTVDMEGARGADAVTIAAVCHTDTSLWNPEHVSFKLLGTPGGTPVCHL
MPYGHIIWAKNVKRSPA

taRAFTIN1d (partial sequence, 82 residues)

MARFLVALLAATLVAVQAGGQLGHAAPATAEVFWRAVLPHSPLPDAVLRLLLKQPAAGVELHTEATSFVR
DPEDRPPFDYRDY

osRAFTIN1 (412 residues)

MARFLLLLVAVAAAAAVLSLGDAAAPSTAEVFWRAVLPELPLDAFLRLLRPDTSFVVGKAEAAGGAART
GFPFDYTDYRGSDSPTTASGLDLAGDFGEPAPFGYDYSAQGEAGGGGAAAAAGEQVLAVDAGFNVDKYV
GARKLRGGSSTAGGENDDEPFGYDYKAPSSSGSGTAASTTARGVGTGATTTVFFHEEAVRVGERLPFYFP
AATTSALGFLPRRVADSIPTAAALPAVLALFGVAPDTAEAAGMRETLRTCEWPTLAGESKFCATSLEA
LVEGAMAALGTRDIAALASTLPRGGAPLQAYAVRAVLPEVAGGFVACHDQAYPYTVYRCHTTGPARAYM
VMEGDGGGDGGEAVTVATVCHTNTSRWNPEHVSFKLLGTPGGSPVCHLMPYGHIVWAKNVKSSTA

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